

SEQUENCE LISTING

- <110> ISHIWATA, TETSUYOSHI
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- <110> SAWADA, SHIGEMASA
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Met Asn His Pro Trp

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846

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35 40 45

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274

120

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aatagttaat	agctgtatta	gccagaaaat	ggtgtaagga	caacaggcta	actaaccctg	180
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ttacataaga	cattctcggt	aagccccctt	tgggtatccc	aaataaggac	tggggtgggt	180
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ttatcttatc	tgaataattt	tgtctgttga	ctattgggat	agttctcctt	cagatgagct	180
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atgttaagca	tcaaggaaaa	taaaacacat	cattgcacat	tacageegea	aaaaac	176
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taaatgccac	acctgatgga	gtcattaggc	actttcctag	tgacaagtgc	ctaggacaga	180
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aggcaacact	ttgctcacaa	tcctacagat	ctacttcacc	tgtaactaca	attttcctga	180
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tcacctagaa	gctctgttta	tttctgagca	accctggggc	ttgtctcata	ggacaggatt	180
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gggagcctga	a ggcatgagaa	a tcacttgaad	c ctgggaggt	g gaggttgcca	a tgagccgaga	180
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<210> 32

<211> 298

<212> DNA

<213> Homo sapiens

<400> 32

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etetetggea gtgeteeetg caaatgtgte ettteaagaa aacaaaacet geaagtgget 180
tgtaatgtae catgaeetta teatgtgaag gacaaatgge tettgtgett attagatage 240
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<210> 33

<211> 291

<212> DNA

<213> Homo sapiens

<400> 33

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agaatgaacc agagccagga cagcgtaaag gctaggttca cagtgagatg aaagaacctg 120
aaaacaagtt taaaactcaa aagaggatta ttctcaagtt atactacagt gaaaaaacat 180
ggaaaaacac aaaaaggaca ggcaataagg cacaggcata catacaaggc aaattgtaac 240
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<210> 34

<211> 230

<212> PRT

<213> Homo sapiens

<400> 34

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Ser Gly Leu Gly Ser Pro His Cys Phe Ser His Gln Asn Gly Glu Arg

Val Glu Arg Tyr Ser Arg Lys Val Phe Val Gly Gly Leu Pro Pro Asp

Ile Asp Glu Asp Glu Ile Thr Ala Ser Phe Arg Arg Phe Gly Pro Leu

Ile Val Asp Trp Pro His Lys Ala Glu Ser Lys Ser Tyr Phe Pro Pro

Lys Gly Tyr Ala Phe Leu Leu Phe Gln Asp Glu Ser Ser Val Gln Ala

Leu Ile Asp Ala Cys Ile Glu Glu Asp Gly Lys Leu Tyr Leu Cys Val

Ser Ser Pro Thr Ile Lys Asp Lys Pro Val Gln Ile Arg Pro Trp Asn

Leu Ser Asp Ser Asp Phe Val Met Asp Gly Ser Gln Pro Leu Asp Pro

Arg Lys Thr Ile Phe Val Gly Gly Val Pro Arg Pro Leu Arg Ala Val

Glu Leu Ala Met Val Met Asp Arg Leu Tyr Gly Gly Val Cys Tyr Ala

Gly Ile Asp Thr Asp Pro Glu Leu Lys Tyr Pro Lys Gly Ala Gly Arg

Val Ala Phe Ser Asn Gln Gln Ser Tyr Ile Ala Ala Ile Ser Ala Arg

Phe Val Gln Leu Gln His Gly Glu Ile Asp Lys Arg Val Ser Leu Ile

Leu His Phe Gly Lys Phe 230 225

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<211> 143

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<213> Homo sapiens

<400> 35

Met Gly Ser Asp Lys Arg Val Ser Arg Thr Glu Arg Ser Gly Arg Tyr 10 5 1

Gly Ser Ile Ile Asp Arg Asp Asp Arg Asp Glu Arg Glu Ser Arg Ser 25 20

Arg Arg Arg Asp Ser Asp Tyr Lys Arg Ser Ser Asp Asp Arg Gly 40 35

Asp Arg Tyr Asp Asp Tyr Arg Asp Tyr Asp Ser Pro Glu Arg Glu Arg 60 55 50

Glu Arg Arg Asn Ser Asp Arg Ser Glu Asp Gly Tyr His Ser Asp Gly 75 70 65

Asp Tyr Gly Glu His Asp Tyr Arg His Asp Ile Ser Asp Glu Arg Glu 90 85

Ser Lys Thr Ile Met Leu Arg Gly Leu Pro Ile Thr Ile Thr Glu Ser 110 105 100

Asp Ile Arg Glu Met Met Glu Ser Phe Glu Gly Pro Gln Pro Ala Asp 120 115

Val Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser 140 135 130

<210> 36

<211> 104

<212> PRT

<213> Homo sapiens

<400> 36

Met Pro His Met Leu Ser Gln Leu Ile Ala Gly Gly Val Ser Thr Ser 15 10 1

Cys Val Thr Ala Leu Gly Glu Glu Thr Gly Ala Trp Phe Pro Val Tyr 20 25 30

Leu Ser His Ala Ser Ser Pro Phe Ala Asp Leu Val Phe Cys Pro Phe 35 40 45

Ala Glu Ile Asn His Ser Gln Glu Tyr Asp Asn Met Arg Gly Pro Val 50 55 60

Ser Pro Pro Asn Lys Gln Phe Asn Leu Gly Val Ile Phe Gly Ile Pro 65 70 75 80

Asn Asn Cys Arg Phe Pro Thr Asp Asn Lys Ile Thr Glu Lys Gln Leu 85 90 95

Leu Gly Asn Val Leu Asn Tyr Pro
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<210> 37

<211> 133

<212> PRT

<213> Homo sapiens

<400> 37

Met Asn His Pro Trp His Val Cys Phe Leu Phe Lys Val Leu Arg Tyr
1 5 10 15

Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 20 25 30

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 35 40 45

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 50 55 60

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys 65 70 75 80

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 85 90 95

Ser Leu Glu Pro Ala Val Ala Glu His Trp Ser Gly Glu Phe Glu Lys 100 105 110 Trp Lys Val Gly Phe Phe His Pro Leu Lys Arg Glu Asp Gly Phe Phe 120 115

Thr Arg Thr Asp Ile 130

<210> 38

<211> 133

<212> PRT

<213> Homo sapiens

<400> 38

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Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 25 20

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 40 35

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 60 55 50

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys 75 70 65

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 90 85

Ser Leu Glu Pro Ala Phe Ala Glu His Trp Ser Gly Glu Phe Glu Lys 110 105 100

Trp Lys Val Gly Phe Phe His Pro Leu Lys Arg Glu Asp Gly Phe Phe 125 120 115

Thr Arg Thr Asp Ile 130

<210> 39

<211> 128

<212> PRT

<213> Homo sapiens

<400> 39

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Glu Lys Leu Leu Leu Ala Thr Gly Leu Asp Gly Ser Tyr Leu Leu Arg

Asp Ser Glu Ser Val Pro Gly Val Tyr Cys Leu Cys Val Leu Tyr His

Gly Tyr Ile Tyr Thr Tyr Arg Val Ser Gln Thr Glu Thr Gly Ser Trp

Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Lys Ile

Lys Asn Leu Ile Ser Ala Phe Gln Lys Pro Asp Gln Gly Ile Val Ile

Pro Leu Gln Tyr Pro Val Glu Lys Lys Ser Ser Ala Arg Ser Thr Gln

Gly Thr Thr Gly Ile Arg Glu Asp Pro Asp Val Cys Leu Lys Ala Pro

<210> 40

<211> 343

<212> PRT

<213> Homo sapiens

<400> 40

Met Asp Ala Pro Lys Ala Gly Tyr Ala Phe Glu Tyr Leu Ile Glu Thr

Leu Asn Asp Ser Ser His Lys Lys Phe Phe Asp Val Ser Lys Leu Gly

Thr Lys Tyr Asp Val Leu Pro Tyr Ser Ile Arg Val Leu Leu Glu Ala

Ala Val Arg Asn Cys Asp Gly Phe Leu Met Lys Lys Glu Asp Val Met

Asn Ile Leu Asp Trp Lys Thr Lys Gln Ser Asn Val Glu Val Pro Phe

Phe Pro Ala Arg Val Leu Leu Gln Asp Phe Thr Gly Ile Pro Ala Met 85 90 95

Val Asp Phe Ala Ala Met Arg Glu Ala Val Lys Thr Leu Gly Gly Asp

Pro Glu Lys Val His Pro Ala Cys Pro Thr Asp Leu Thr Val Asp His 115 120 125

Ser Leu Gln Ile Asp Phe Ser Lys Cys Ala Ile Gln Asn Ala Pro Asn 130 135 140

Pro Gly Gly Gly Asp Leu Gln Lys Ala Gly Lys Leu Ser Pro Leu Lys 145 150 150

Val Gln Pro Lys Lys Leu Pro Cys Arg Gly Gln Thr Thr Cys Arg Gly
165 170 175

Ser Cys Asp Ser Gly Glu Leu Gly Arg Asn Ser Gly Thr Phe Ser Ser 180 185 190

Gln Ile Glu Asn Thr Pro Ile Leu Cys Pro Phe His Leu Gln Pro Val 195 200 205

Pro Glu Pro Glu Thr Val Leu Lys Asn Gln Glu Val Glu Phe Gly Arg 210 215 220

Asn Arg Glu Arg Leu Gln Phe Phe Lys Trp Ser Ser Arg Val Leu Lys 225 230 230

Asn Val Ala Val Ile Pro Pro Gly Thr Gly Met Ala His Gln Ile Asn 245 250 255

Leu Glu Tyr Leu Ser Arg Val Val Phe Glu Glu Lys Asp Leu Leu Phe 260 265 270

Pro Asp Ser Val Val Gly Thr Asp Ser His Ile Thr Met Val Asn Gly 275 280 285

Leu Gly Ile Leu Gly Trp Gly Val Gly Gly Ile Glu Thr Glu Ala Val 290 295 300

Met Leu Gly Leu Pro Val Ser Leu Thr Leu Pro Glu Val Val Gly Cys 305 310 315 320

Glu Leu Thr Gly Ser Ser Asn Pro Phe Val Thr Ser Ile Asp Val Val

325 330 335

Leu Gly Ile Thr Lys Val Ser 340

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<211> 305

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cataatttaa acaaaatcaa ctaagatgat ccaagttcca cacaactgca cttcaatatt 180
caagtcggtg tgaagatgcc tgactactgc gtcacaagat tctgagctgt cgtaaaaagc 240
ctggctcgtg gtttctattt atagtgtaca catgttgggt tataatcaca aacctggaac 300
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<210> 42

<211> 256

<212> DNA

<213> Homo sapiens

<400> 42

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ttgggaagca gctttacaaa tgtgacttga cttggggatc ttcttgatac tttgccatgg 180

caaggaacaa gccgcctgaa	ctaaatgcca	ctccatttga	ttccacgctt	aaagtaacca	240
tgcaaccgac tatagt					256
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tgagetteet geettaaate	atacccacag	tgaatggcgt	ccctttatca	ccgctaatga	180
ctctgacatc tctctccact	cacatgtgag	cctcctcagc	tctcganaaa	caagtengte	240
tcgg					244
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tactctccaa tggtgatgaa	gggagatgtc	tgggggaaat	ccagcaggat	gttgatttag	120
tatgtacaca gtgagaggat	acttgtagag	aacctagaat	cttctctgaa	tgtgacgggc	180
cctcagagat aattgttaac	agataagtgg	atgattaaat	acacttcctc	cagtaggcta	240

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258
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<211> 26
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<210> 46
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 46
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gttattatac tatcaagtaa cccaac
<210> 47
<211> 25
<212> DNA
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gtggatctgg atttttgtca tatgt
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<211> 25
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<213> Artificial Sequence

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gacaatgagt aagaagaaag aggg	24

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ggtacccagt ttcaaattaa catgg
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<210> 55
<211> 24
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gaccaggact gaacagaggt ga
<210> 57
<211> 25
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<400> 57
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gcttatagac catgtttgta gtagg
<210> 58
<211> 25
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<210> 59 <211> 22

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gccacgggtt tcccatatcg aa
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<400> 61
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gttctgctct cagcagattg gtta
<210> 62
<211> 24
<212> DNA
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<400> 67 gagacagcat tcagatatag acgg	24
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<400> 69 gatggcctgt gtgaacagat taat	24

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<211> 24
<212> DNA
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<210> 71
<211> 24
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gatccccaca atttcttgtg attg
<210> 72
<211> 25
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<211> 23
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<210> 78
<211> 22
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<223> Description of Artificial Sequence: Synthetic DNA
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                                                                         22
<210> 79
<211> 24
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<211> 25
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<223> Description of Artificial Sequence: Synthetic DNA
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<400> 81
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<210> 82
<211> 24
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<400> 82
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<210> 83
<211> 26
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<210> 84
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<211> 24

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<210> 85
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<400> 85
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gaggtagggc ttcccttcgc ta
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<211> 25
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<223> Description of Artificial Sequence: Synthetic DNA
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gcataacaag tgacagggtt agtta
<210> 87
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 87
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22
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<210> 88
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
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<210> 89
<211> 24
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
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<210> 90
<211> 25
<212> DNA
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<220>
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<400> 90
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<211> 24 <212> DNA

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<210> 93 <211> 24 <212> DNA <213> Artificial Sequence	
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<400> 93	
gacagcaacc taataacagc tgtc	24
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<212> DNA <213> Artificial Sequence	
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<210> 95
<211> 22
<212> DNA
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 95
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gaggggactt ccaagagtct ct
<210> 96
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 96
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gtcttcagga aaattgtagt tacag
<210> 97
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 97
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gttacaaaca cacacgaagt tcct
<210> 98
<211> 22
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 98
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gacttectaa ggeacaetea ge
<210> 99
<211> 24
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 99
                                                                         24
gtttaactac ctctcaggtc atga
<210> 100
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 100
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gtcgccaagg ctgtagtgca at
<210> 101
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 101
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gaaataggta tcccttgatg tcga
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<210> 102
<211> 24
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA
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gaccaagaat tcagttcatc agtt
<210> 103
<211> 22
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 103
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gaatgaacca gagccaggac ag
<210> 104
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 104
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gccttgtatg tatgcctgtg cc
<210> 105
<211> 21
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
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<400> 105
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aagagtccac caggccatgg a
<210> 106
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 106
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taccttgtgt acttctagct gag
<210> 107
<211> 17
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 107
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gtttttttt ttttta
<210> 108
<211> 17
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<213> Artificial Sequence
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gtttttttt tttttg
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<210> 109 <211> 17

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<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 109
gtttttttt tttttc
                                                                        17
<210> 110
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<223> Description of Artificial Sequence: Synthetic DNA
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cagagtgatg gatatcaa
<210> 111
<211> 22
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 111
                                                                        22
atgaaagtgc cagtgtgcca tg
<210> 112
<211> 22
<212> DNA
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<220>
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<400> 112
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26

cccatcacca tcttccagga gc

<210> 116

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<210> 113
<211> 26
<212> DNA
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ttcaccacct tcttgatgtc atcata
<210> 114
<211> 15
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<220>
<223> Description of Artificial Sequence: Synthetic Peptide
<400> 114
Cys Pro Leu Lys Arg Glu Asp Gly Phe Phe Thr Arg Thr Asp Ile
                                      10
                                                           15
  1
<210> 115
<211> 16
<212> PRT
<213> Artificial Sequence
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<221> MOD RES
<222> (16)
<223> AMIDATION, GluAmide
<400> 115
Cys Ser Phe Leu Glu Lys Phe Asn Lys Ser Lys Arg Glu Arg Leu Xaa
                                                           15
  1
                  5
                                      10
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<211> 15
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<220>
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<222> (15)
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<400> 116
Cys Ala Glu His Trp Ser Gly Glu Phe Glu Lys Trp Lys Val Xaa
                                                           15
                                      10
<210> 117
<211> 16
<212> PRT
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<220>
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<400> 117
Cys Glu Ile Asp Lys Arg Val Ser Leu Ile Leu His Phe Gly Lys Phe
                                      10
<210> 118
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic Peptide
<400> 118
Cys Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser
                                                           15
  1
                  5
                                      10
<210> 119
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<220>
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<400> 119
Cys Thr Ser Ile Asp Val Val Leu Gly Ile Thr Lys Val Ser
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                  5
<210> 120
<211> 16
<212> PRT
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<222> (16)
<223> AMIDATION, LysAmide
<400> 120
Cys Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Xaa
                                                           15
                                      10
  1
<210> 121
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic Peptide
<400> 121
Cys Lys Ile Thr Glu Lys Gln Leu Leu Gly Asn Val Leu Asn Tyr Pro
                                                           15
                                      10
```

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